



L Number	Hits	Search Text	DB	Time stamp
1	13512	coli and (swine or pig or cattle)	USPAT;	2003/09/30 11:06
			US-PGPUB;	
			EPO;	
			DERWENT	
2	4944	coli and (swine or pig or cattle) and	USPAT;	2003/09/30 11:06
		(intestine or intestinal)	US-PGPUB;	
			EPO;	
			DERWENT	
3	57	coli and (swine or pig or cattle) and	USPAT;	2003/09/30 11:07
		(intestine or intestinal) and (FUT? or	US-PGPUB;	
1		fucosyltransferase)	EPO;	
			DERWENT	
4	2	(coli and (swine or pig or cattle) and	USPAT;	2003/09/30 11:07
·		(intestine or intestinal) and (FUT? or	US-PGPUB;	
		fucosyltransferase)).clm.	EPO;	1
			DERWENT	
5	10		USPAT;	2003/09/30 11:08
		(intestine or intestinal) and (FUT? or	US-PGPUB;	
		fucosyltransferase) and (alteration or	EPO;	
		mutation or polymorphism).clm.	DERWENT	

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Leu Val Cys Val Leu Ala Ala Ile Phe Phe Leu Asn Val Tyr Gln Asp
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ctc ttt tac agt ggc tta gac ctg ctg gcc ctg tgt cca gac cat aac
                                                                    146
Leu Phe Tyr Ser Gly Leu Asp Leu Leu Ala Leu Cys Pro Asp His Asn
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Val Val Ser Ser Pro Val Ala Ile Phe Cys Leu Ala Gly Thr Pro Val
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cag Gln 95	tat Tyr	gcc Ala	acg Thr	ctg Leu	ctg Leu 100	Ala	ctg Leu	gcg Ala	cag Gln	ctc Leu 105	Asn	ggc Gly	cgc Arg	cag Gln	gcc Ala 110	338
ttc Phe	atc Ile	cag Gln	cct Pro	gcc Ala 115	Met	cac His	gcc Ala	gtc Val	ctg Leu 120	gcc Ala	ccc	gtg Val	ttc Phe	cgc Arg 125	atc Ile	386
acg Thr	ctg Leu	cct Pro	gtc Val 130	ctg Leu	gcg Ala	ccc Pro	gag Glu	gta Val 135	gac Asp	agg Arg	cac His	gct Ala	cct Pro 140	tgg Trp	cgg	434
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gac Asp	tat Tyr	ctg Leu 225	cgt Arg	gtg Val	atg Met	ccc Pro	aag Lys 230	cgc Arg	tgg Trp	aag Lys	Gly 999	gtg Val 235	gtg Val	ggt Gly	gac Asp	722
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Pro Val Leu Ala Pro Glu Val Asp Arg His Ala Pro Trp Arg Glu Leu 135 130 Glu Leu His Asp Trp Met Ser Glu Asp Tyr Ala His Leu Lys Glu Pro 155 150 Trp Leu Lys Leu Thr Gly Phe Pro Cys Ser Trp Thr Phe Phe His His 165 Leu Arg Glu Gln Ile Arg Ser Glu Phe Thr Leu His Asp His Leu Arg 185 Gln Glu Ala Gln Gly Val Leu Ser Gln Phe Arg Leu Pro Arg Thr Gly Asp Arg Pro Ser Thr Phe Val Gly Val His Val Arg Arg Gly Asp Tyr Leu Arg Val Met Pro Lys Arg Trp Lys Gly Val Val Gly Asp Gly Arg 235 Tyr Leu Gln Gln Ala Met Asp Trp Phe Arg Ala Arg Tyr Glu Ala Pro 250 Val Phe Val Val Thr Ser Asn Gly Met Glu Trp Cys Arg Lys Asn Ile Asp Thr Ser Arg Gly Asp Val Ile Phe Ala Gly Asp Gly Arg Glu Ala 280 Ala Pro Ala Arg Asp Phe Ala Leu Leu Val Gln Cys Asn His Thr Ile 295 290 Met Thr Ile Gly Thr Phe Gly Phe Trp Ala Ala Tyr Leu Ala Gly Gly 315 Asp Thr Ile Tyr Leu Ala Asn Phe Thr Leu Pro Thr Ser Ser Phe Leu 330 325 Lys Ile Phe Lys Pro Glu Ala Ala Phe Leu Pro Glu Trp Val Gly Ile 345 Asn Ala Asp Leu Ser Pro Leu Gln Met Leu Ala Gly Pro

360